

# What kind of drugs should we develop

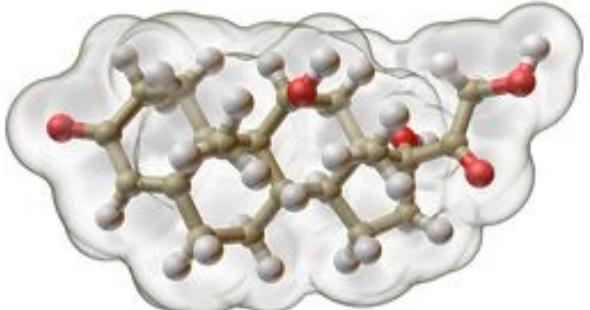
*Mathematical and Computational Biology in Drug Discovery  
(MCBDD) Module III*

*Dr. Jitao David Zhang  
April 2021*

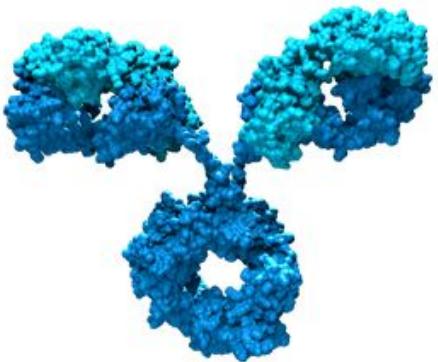
# Overview

- Essentials of modalities
  - Small molecules: classical, protein degrader, RNA modulator
  - Large molecules: classical, DUTA-Fabs, protein design
  - Antisense oligonucleotides: siRNA, shRNA, ASO
  - Gene and cell therapy
- Three case studies:
  - Success stories:
    - [Small molecules] SMA (Evrysdi/Risdiplam and Nusinersen)
    - [Antisense] patisiran ([KEGG DRUG](#)) and givosiran ([DrugBank](#), [structure available at EMA](#))
    - [Offline read] mRNA vaccine (MIT Technology Review)
    - Turning failure into successes: [Multispecific drugs] Thalidomide, PROTAC, degraders
    - [Antibody] Cancer immunotherapy (CTLA4, PD1)
    - [Gene and Cell therapy] CAR-T
  - Challenges
    - [Antisense] HTT (Tominersen)
    - Difference between genetic and enzymatic inhibition

# A zoo of modalities



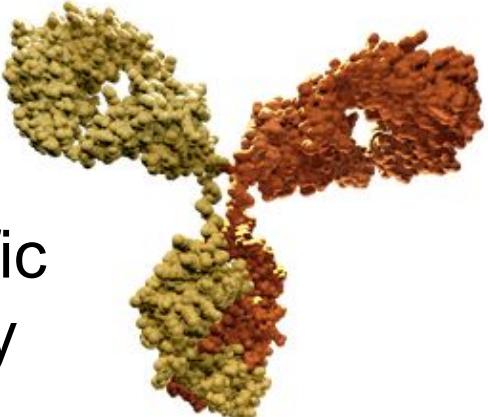
Small molecule



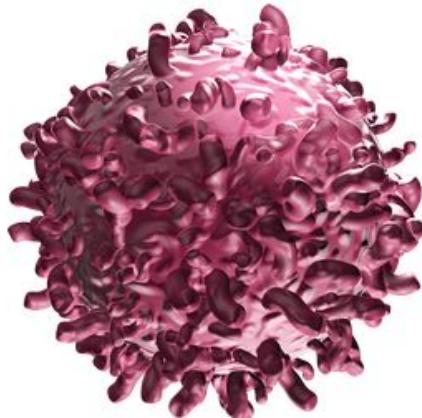
Monoclonal antibody



RNA inference

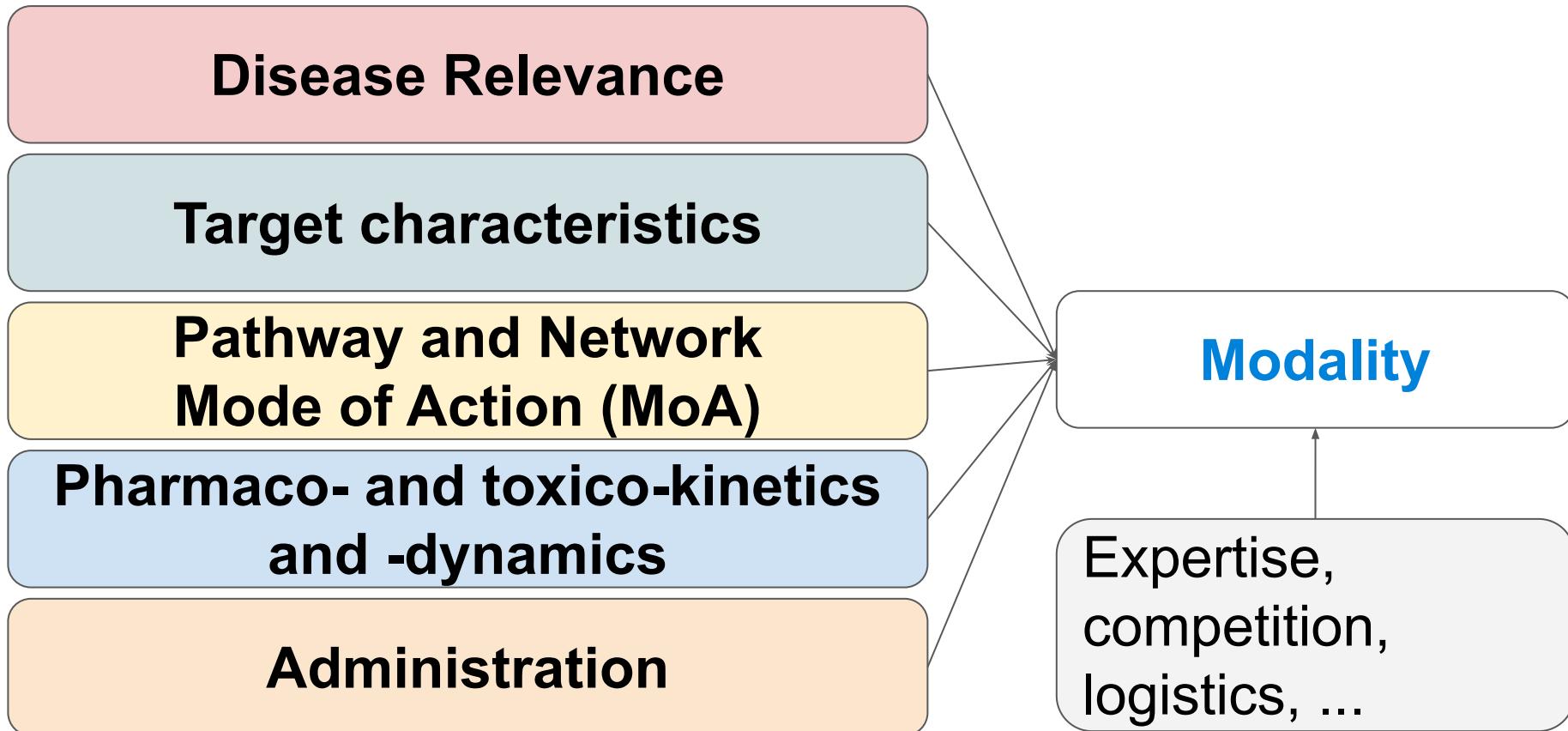


Bispecific  
antibody



Chimeric  
Antigen  
Receptor  
(CAR) T-cells

# Criteria to choose a modality

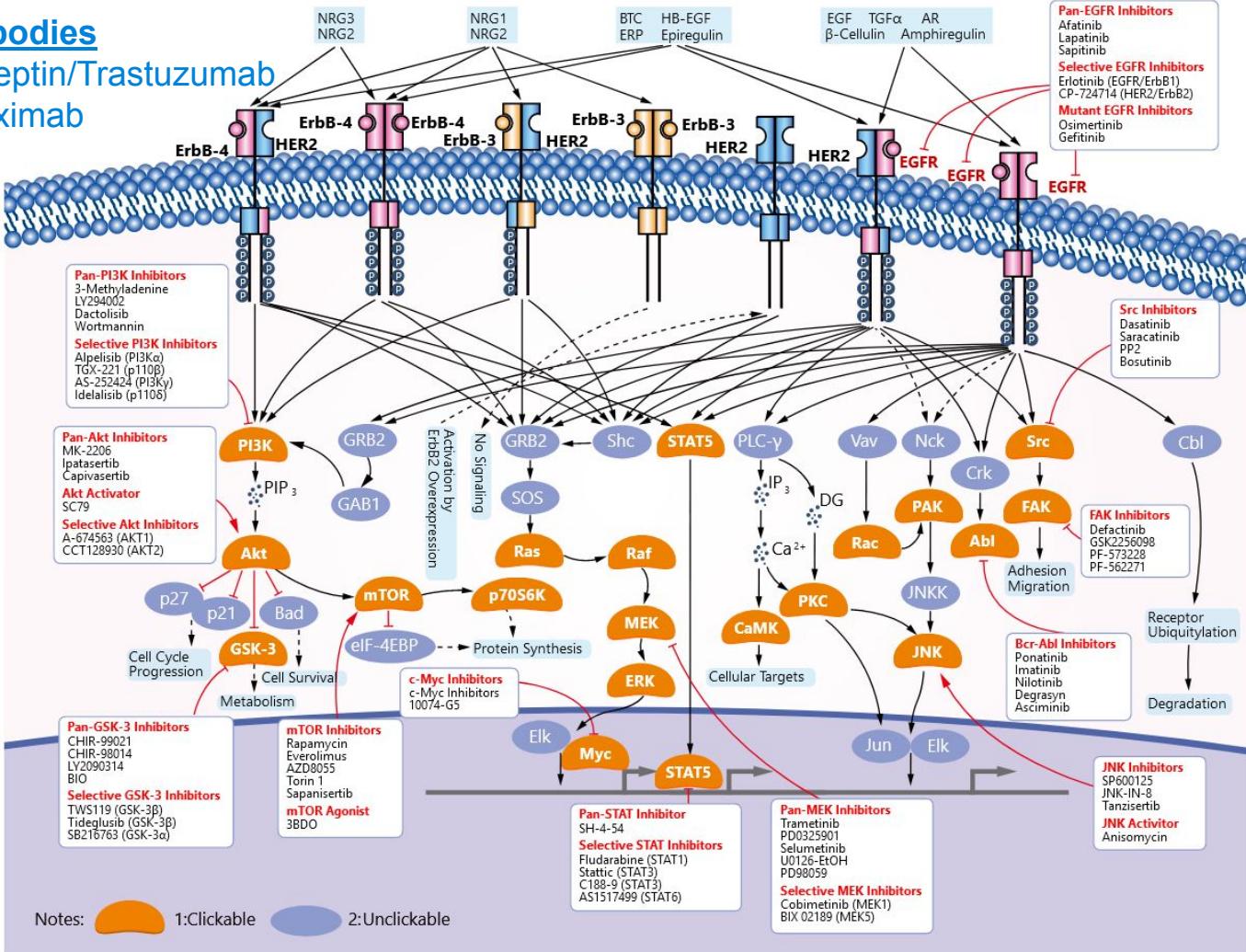


# Multiple modalities can target the same biological process

An example: the epidermal growth factor receptor (EGFR) pathway

## Antibodies

Herceptin/Trastuzumab  
Cetuximab



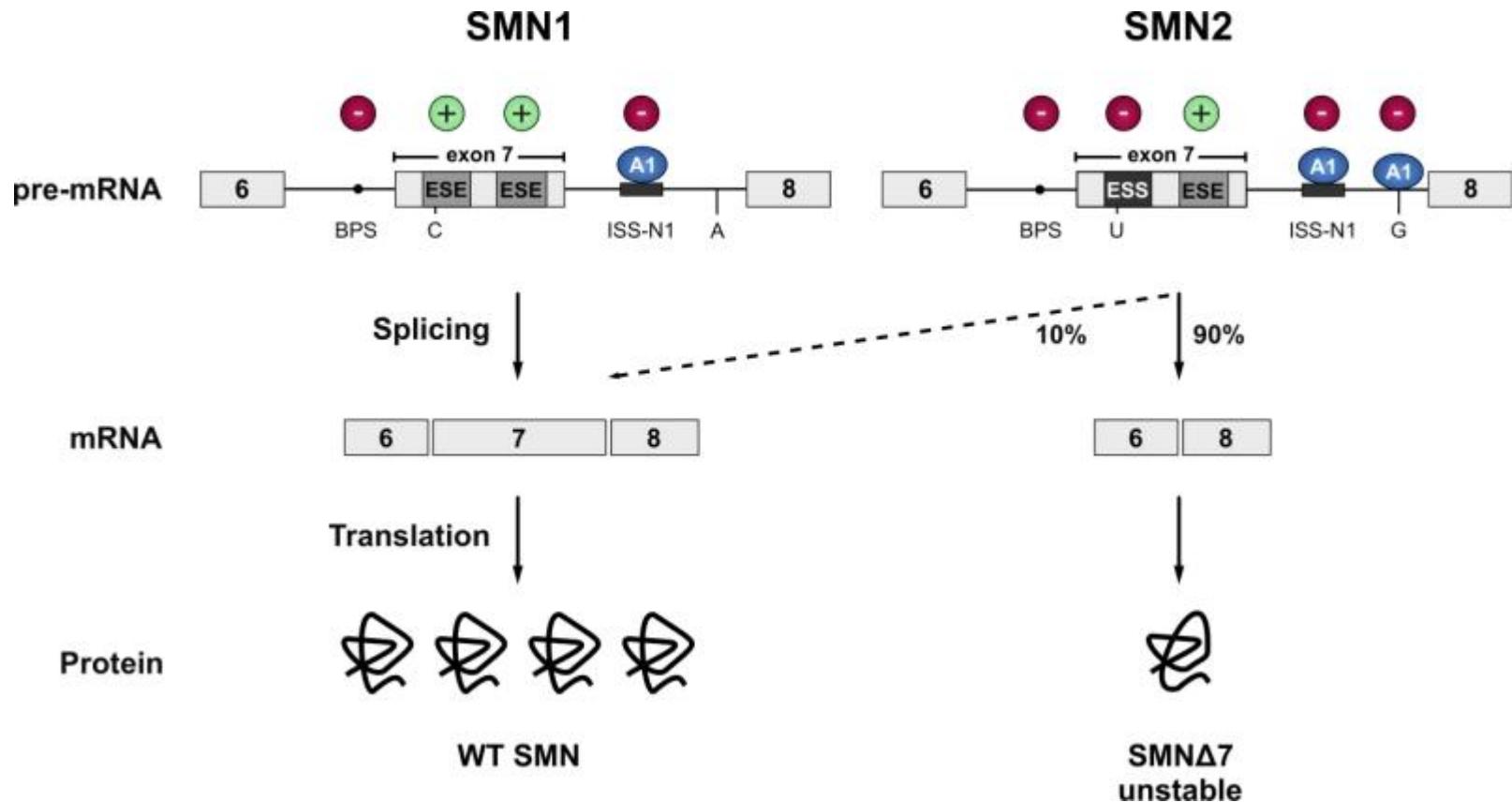
# Characteristics of therapeutic modalities

Modality	Cause of disease at the protein level		Molecular target	Protein target localization			Delivery
	Reduction or loss of function	Excessive or detrimental function		Extracellular	Plasma membrane	Intracellular	
Small molecule	●	●	DNA → RNA → Protein	●	●	●	Oral Injection Inhaled
Protein replacement	○			○	○	○	○
Antibody		●		●	●		●
Oligonucleotide therapy	○	○	○	○	○	○	○
Cell and gene therapy*	●		●	●	●		●

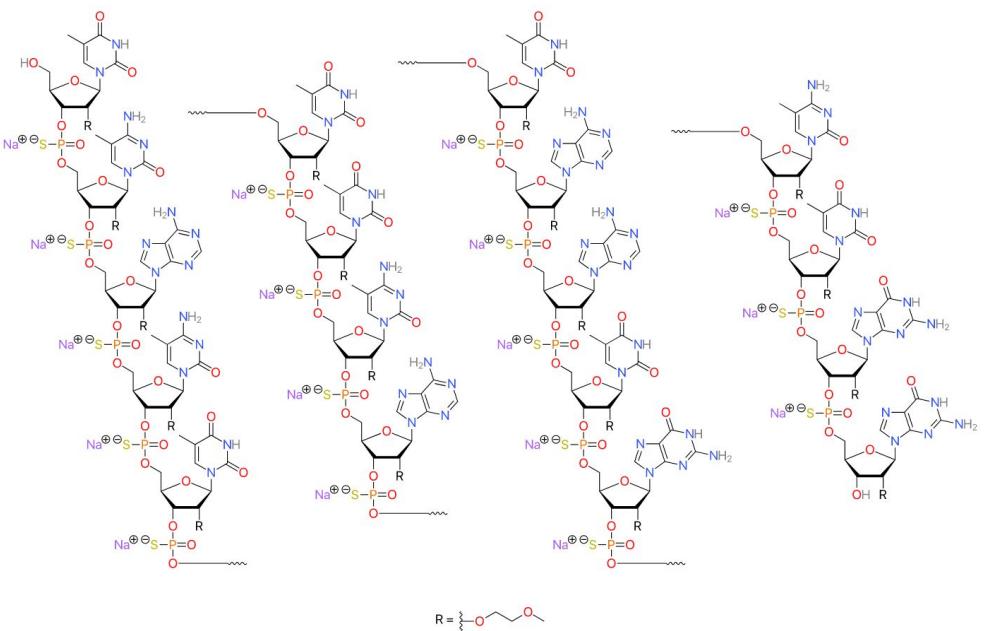
# Facts about Spinal Muscular Atrophy (SMA)

- SMA is caused by a defect in a gene called *SMN1*. People with SMA have reduced levels of the SMN protein.
- When SMN protein levels are reduced, motor neurons are unable to send signals to the muscles, causing them to become smaller and weaker over time
- Depending on the severity, or type of SMA, people with the disease will have difficulties moving, eating, and in some cases breathing, making them increasingly dependent on parents and caregivers
- A short movie: <https://www.nejm.org/doi/full/10.1056/NEJMoa2009965>

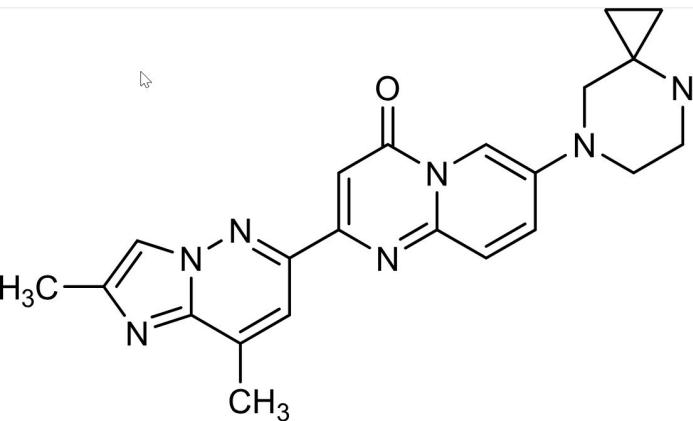
# The molecular mechanism of SMA



# Two Drugs, One Disease

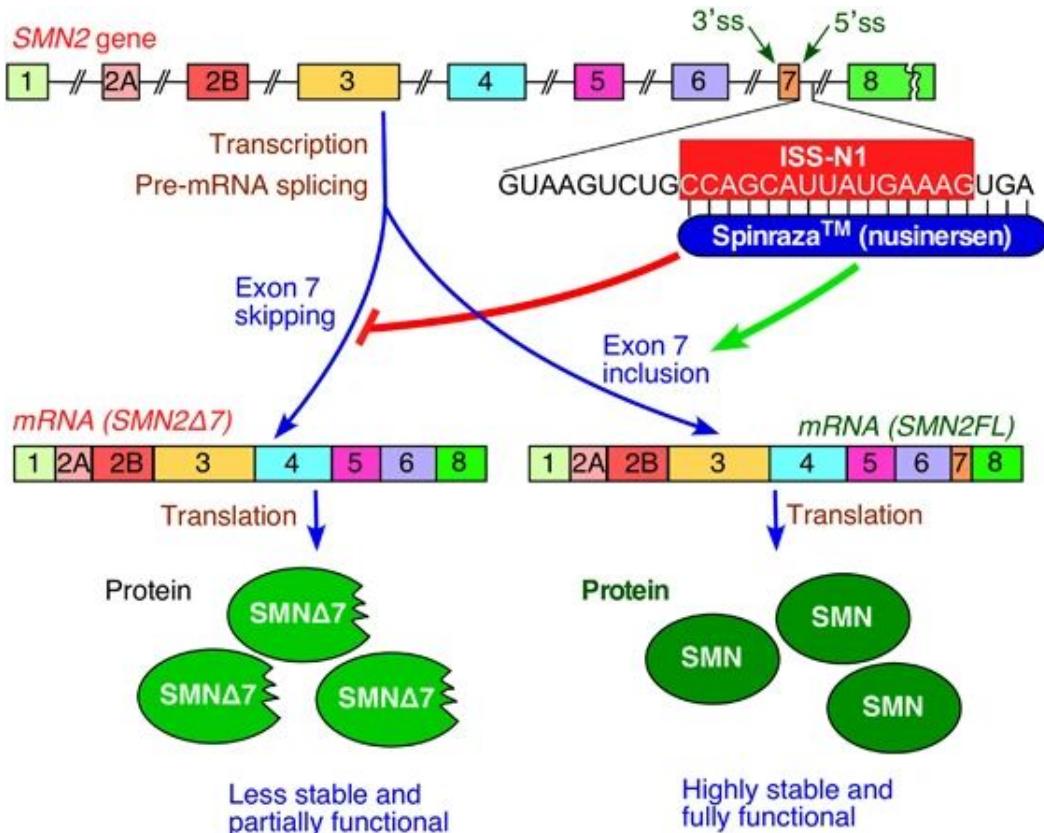


Nusinersen sodium/ Spinraza  
(CHEMBL3833342)



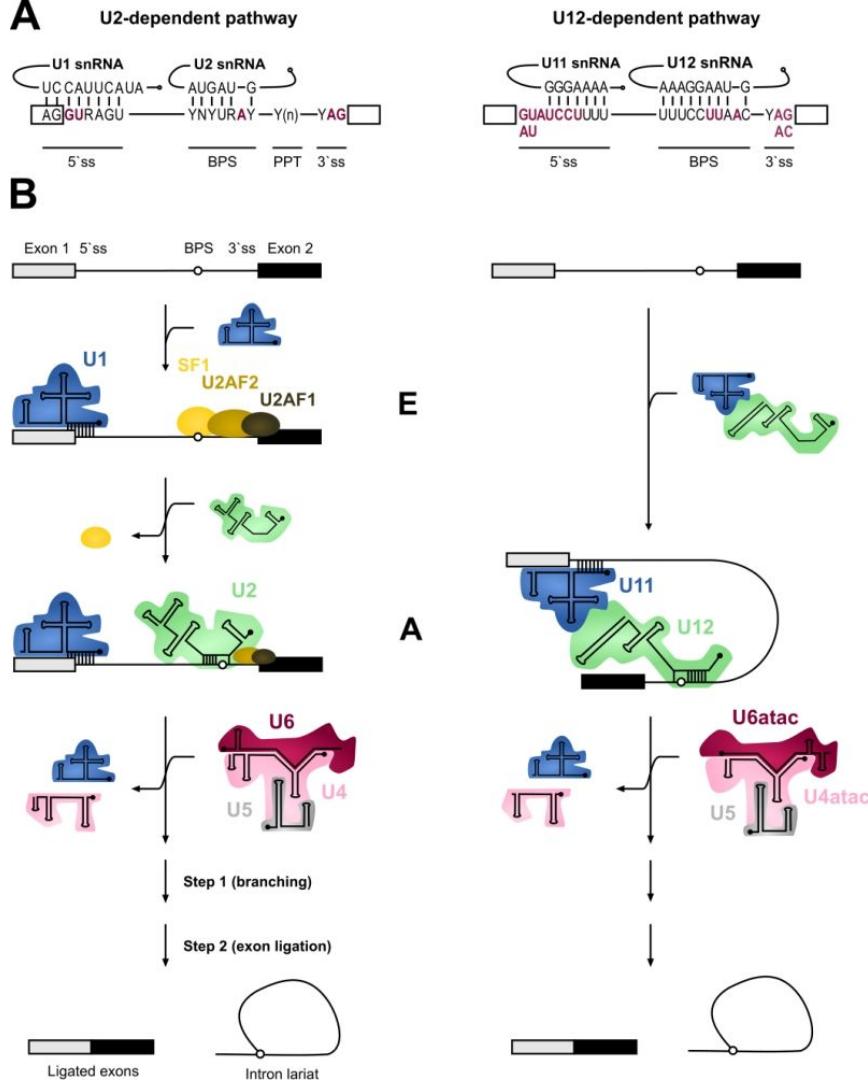
Risdiplam/ Evrysdi/  
(CHEMBL4297528)

# How Spinraza (nusinersen) works

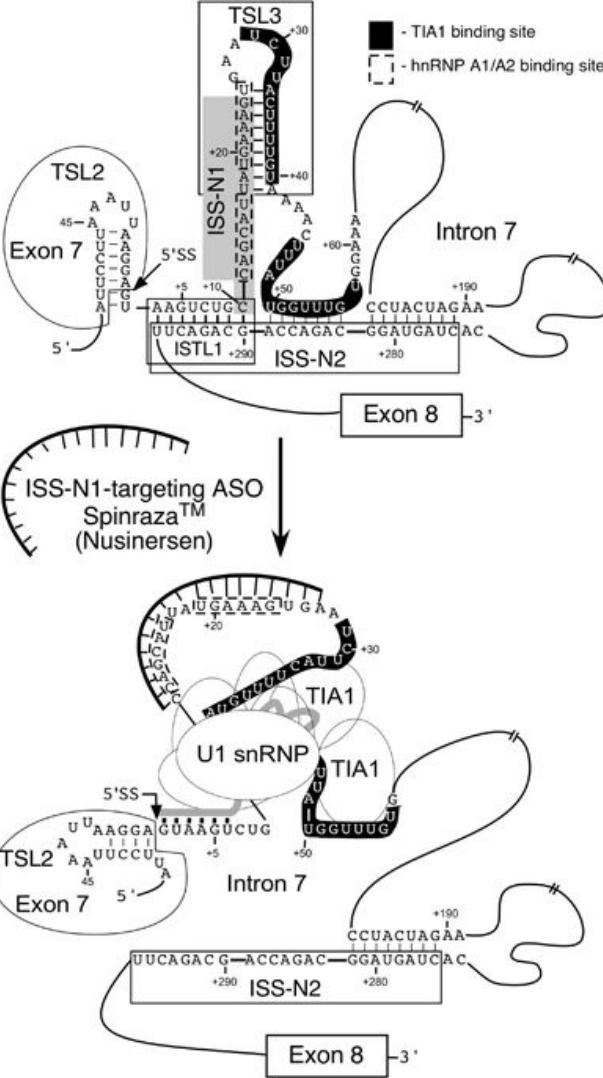


# The splicing machinery

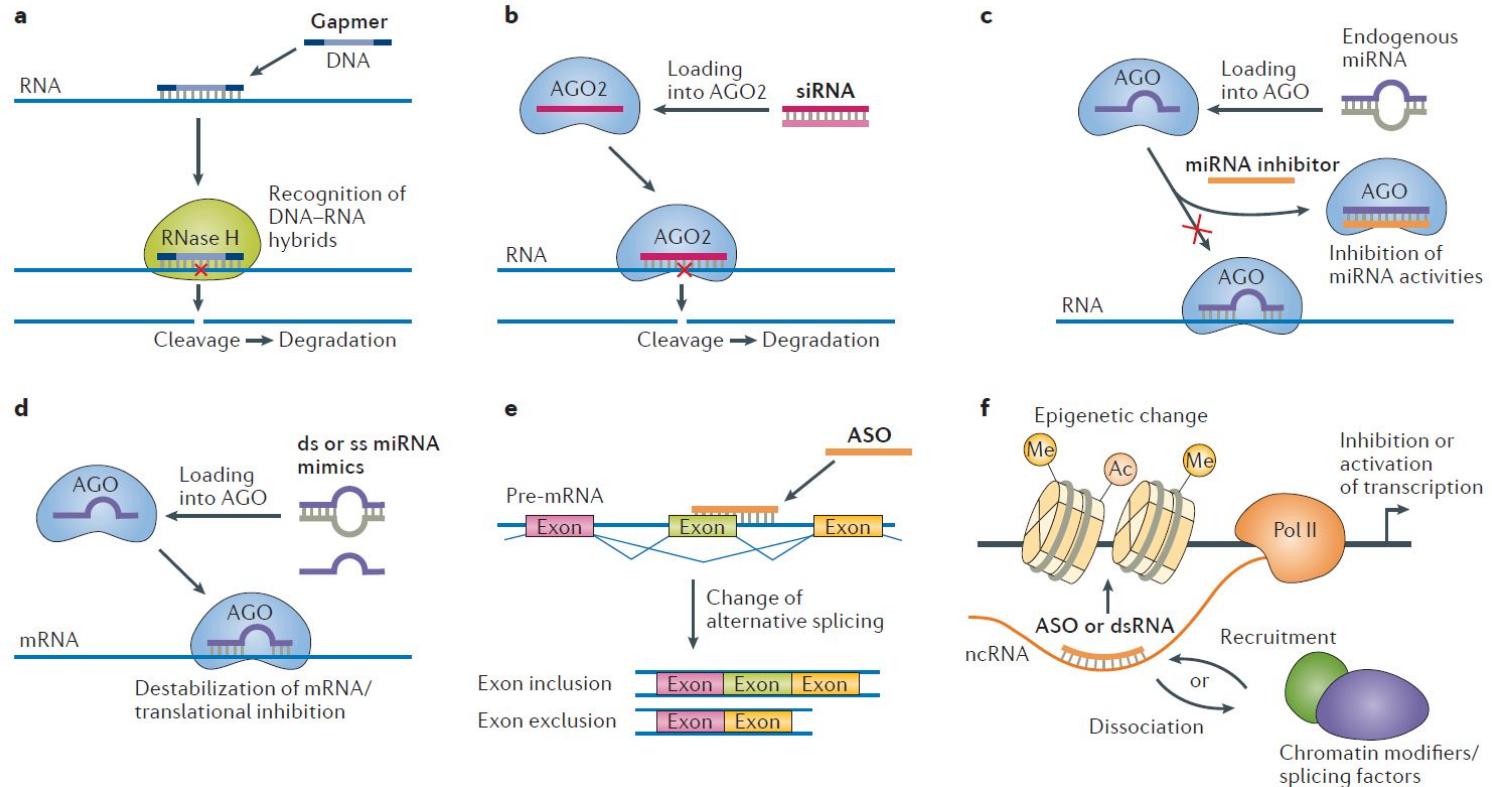
Spliceosome	snRNAs	Core associated proteins
<b>Major</b>	U1	Sm proteins <sup>+</sup> , U1-A , U1-C, U1-70K
	U2	Sm proteins <sup>+</sup> , <b>12S<sup>#</sup></b> : U2-A', U2-B'', 17S <sup>#</sup> : SF3a and SF3b complexes, hPrp43
	U5*	Sm proteins <sup>+</sup> , <b>20S<sup>#</sup></b> : 52K, 40K, hPrp8, hBrr2, Snu114, hPrp6, hPrp28, hDib1
	U4/U6	Sm proteins <sup>+</sup> , LSm proteins2-8, <b>13S<sup>#</sup></b> : CypH, 15.5K, hPrp3, hPrp31, hPrp4
<b>Minor</b>	U11/U12	Sm proteins <sup>+</sup> , <b>18S<sup>#</sup></b> : SF3b complex, 20K ( <i>ZMAT5</i> ), 25K ( <i>SNRNP25</i> ), 31K ( <i>ZCRB1</i> ) 35K ( <i>SNRNP35</i> ), 48K ( <i>SNRNP48</i> ), 59K ( <i>PDCD7</i> ) 65K ( <i>RNPC3</i> ), Urp ( <i>ZRSR2</i> )
	U4atac/U6atac	Share proteins with U4/U6 snRNAs of the major spliceosome



# How Spinaraza (nusinersen) works, base by base

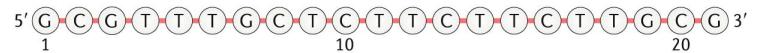


# Regulating RNA levels or splicing with ASOs and duplex RNAs

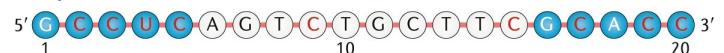


# Chemistry of oligonucleotides evolves with time

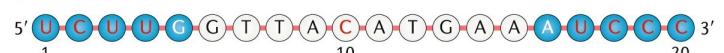
a Fomivirsen



b Mipomersen



c Inotersen



d Eteplirsen



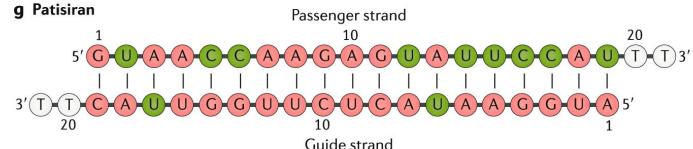
e Golodirsen



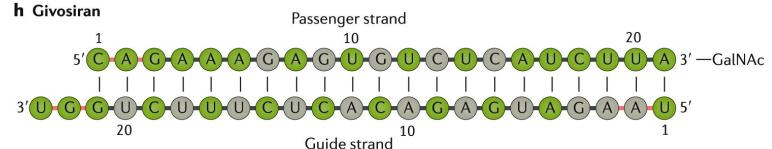
f Nusinersen



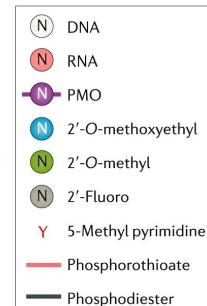
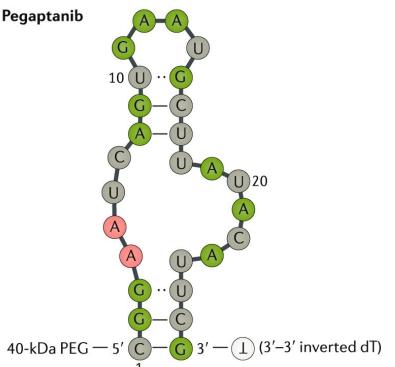
g Patisiran



h Givosiran

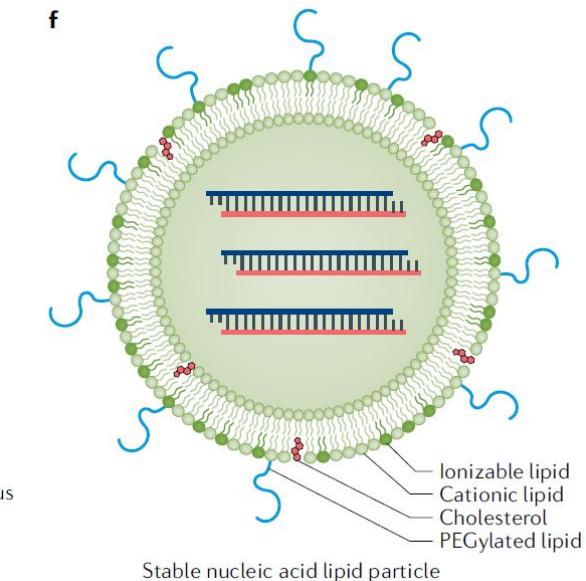
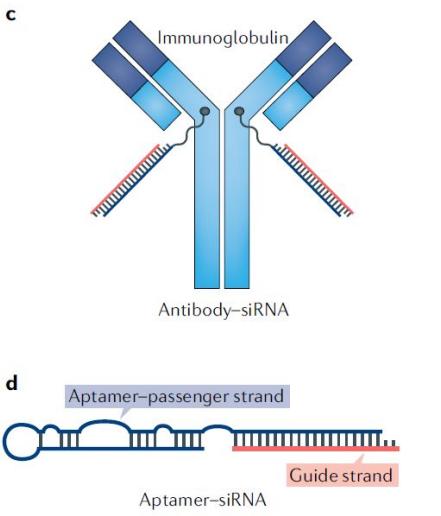
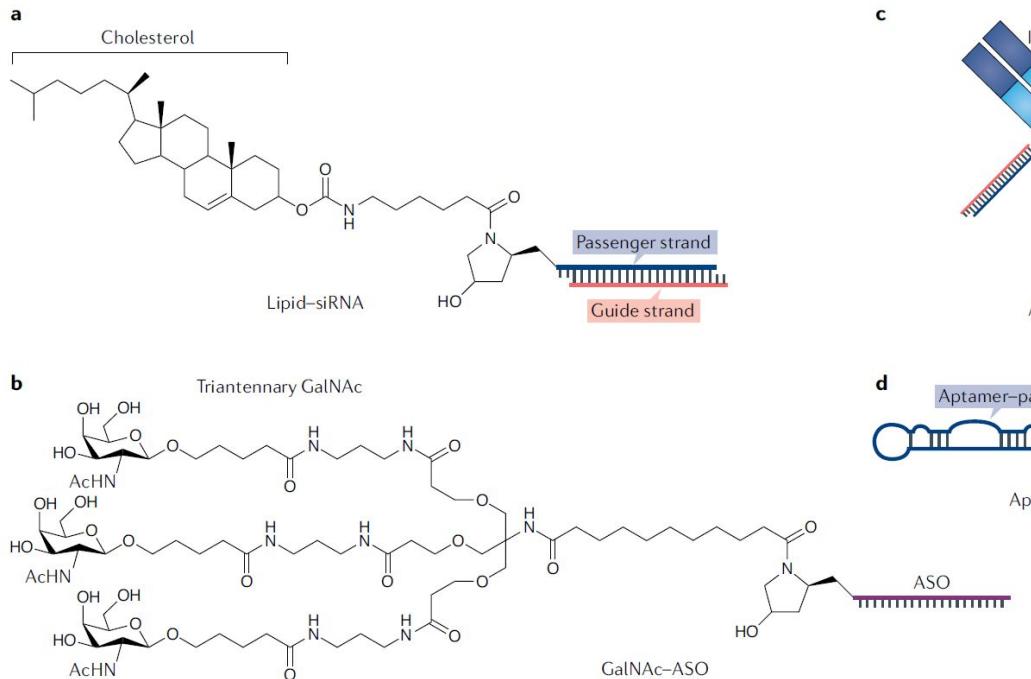


i Pegaptanib

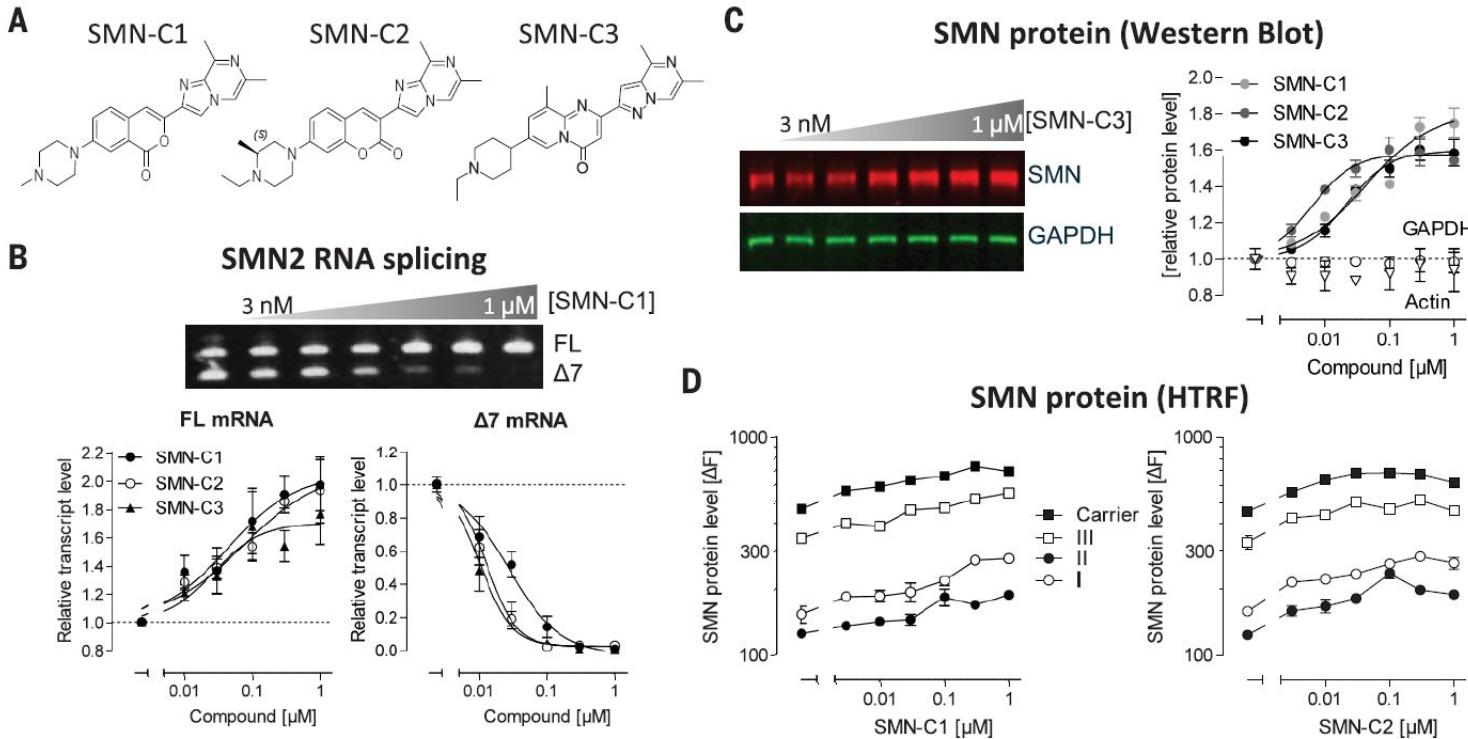


PMO=phosphorodiamidate morpholino oligomer

# Delivery systems of antisense oligonucleotides



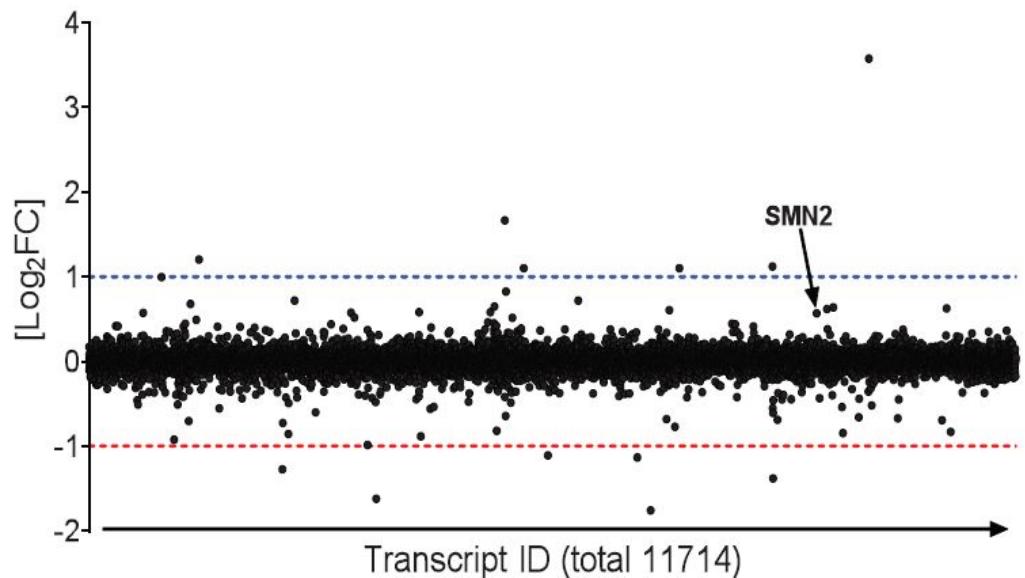
# Small molecules as RNA splicing modifiers



# RNA sequencing confirms the specificity of SMN-C3

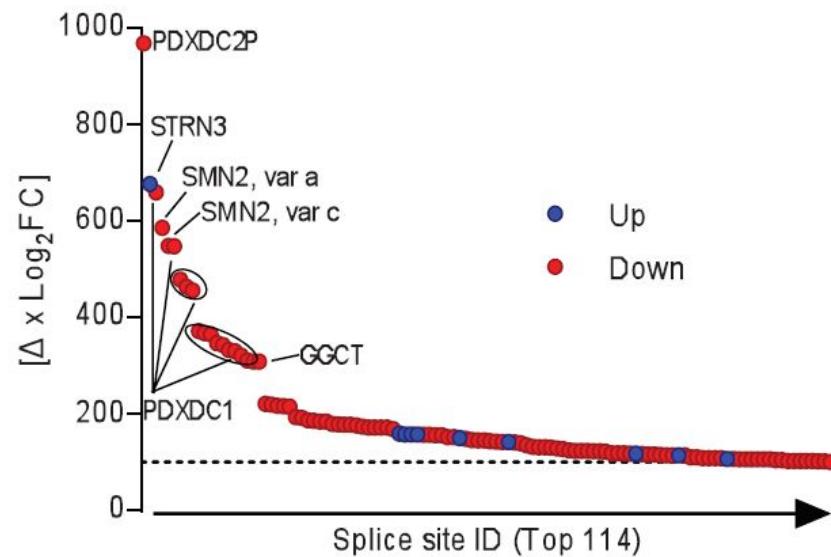
A

Transcriptional changes by SMN-C3



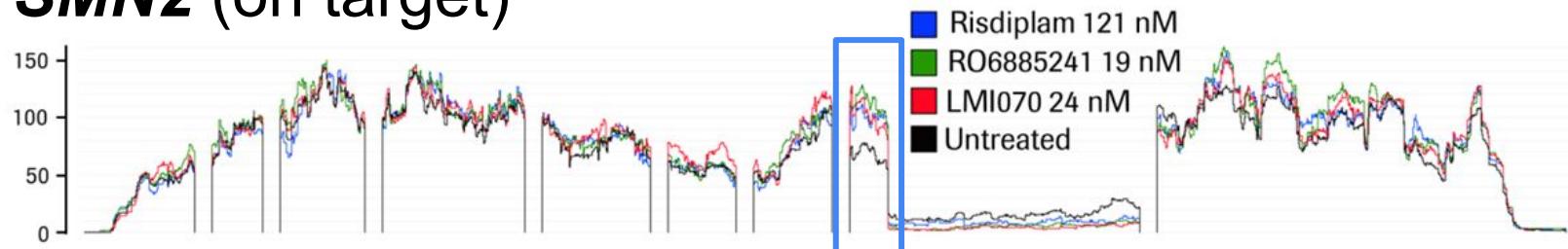
B

Splicing regulation by SMN-C3

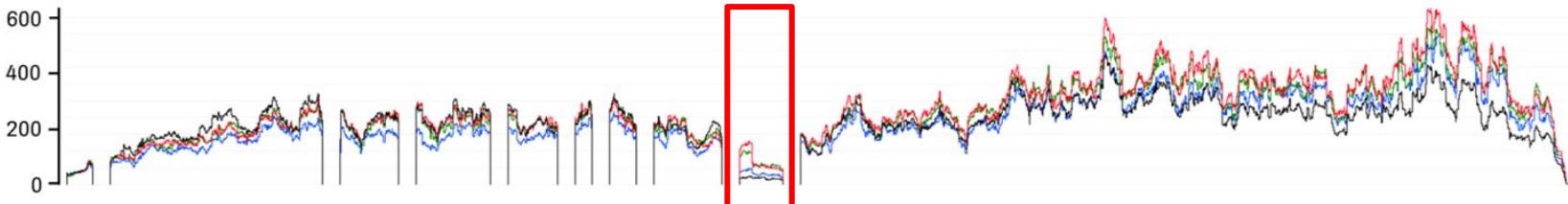


# RNA sequencing confirms the specificity of SMN-C3

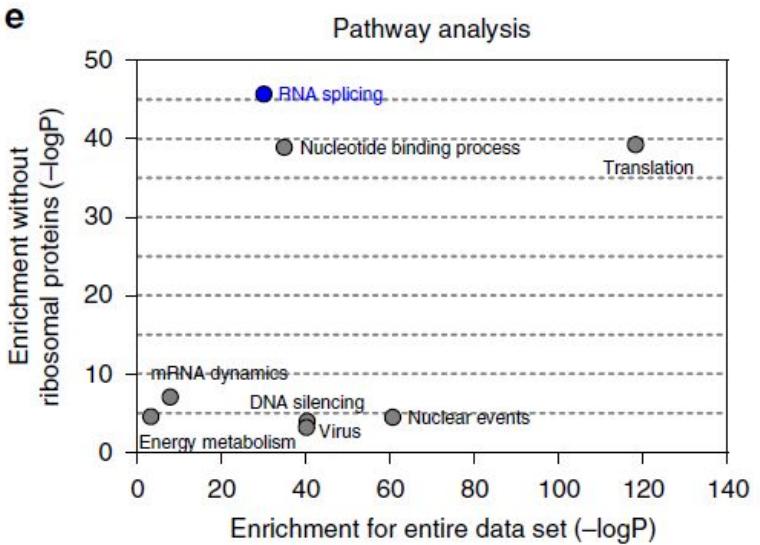
***SMN2* (on target)**



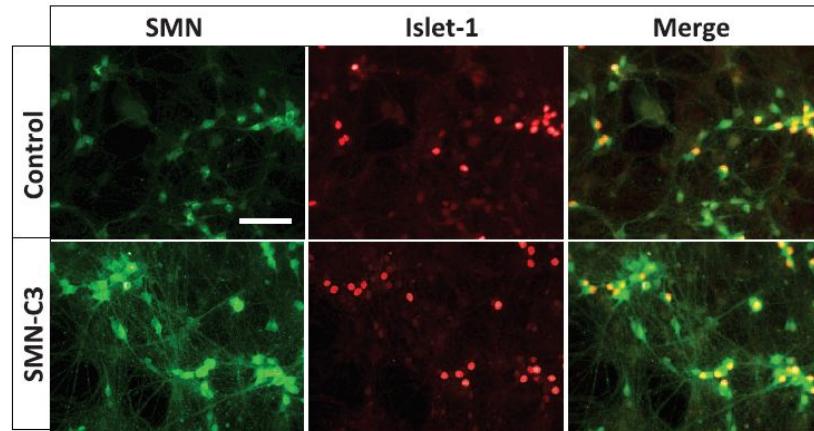
***FOXM2* (off target)**



# Gene-enrichment analysis confirms specific regulation of RNA splicing



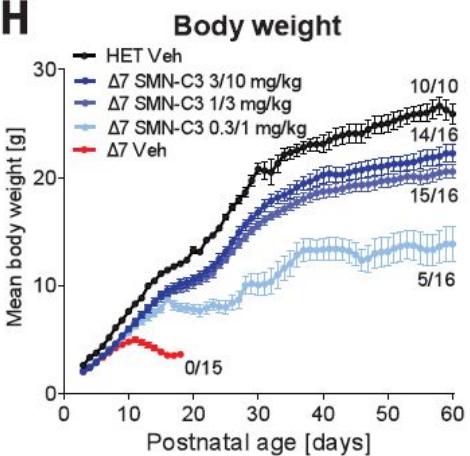
# Experiments *in vitro* and *in vivo* support efficacy profiles of SMN-C3



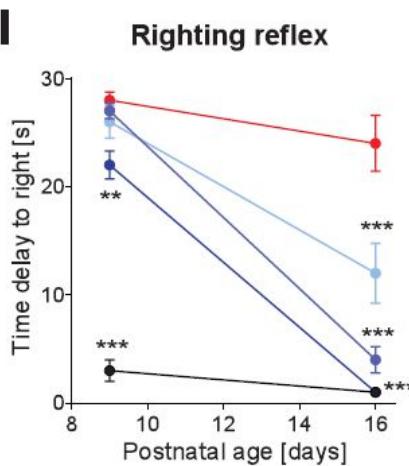
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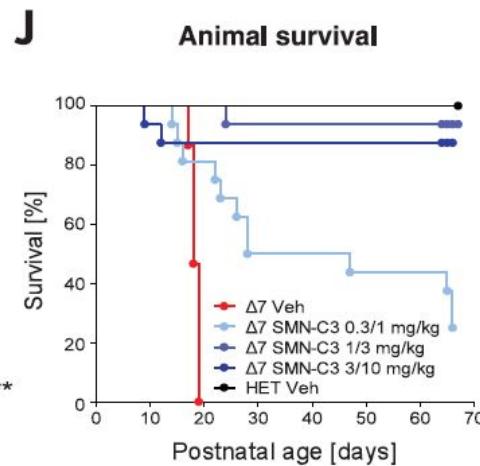
H



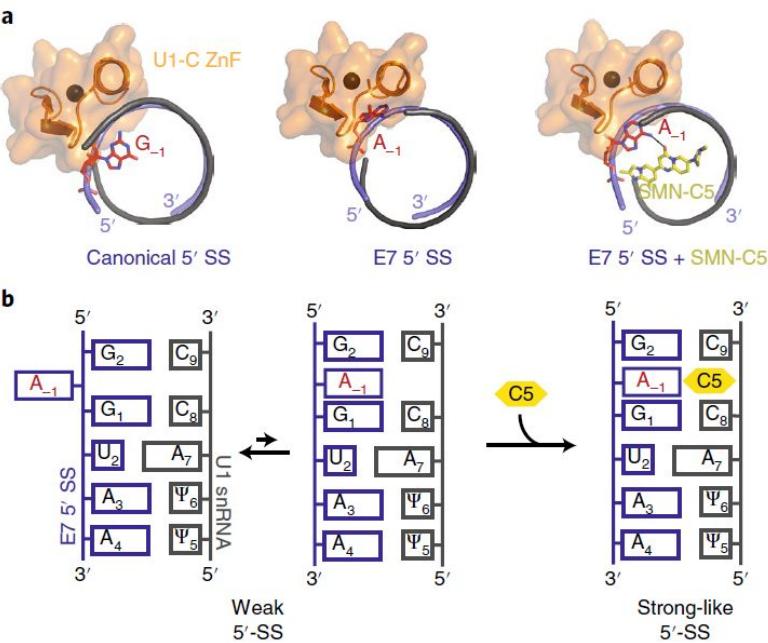
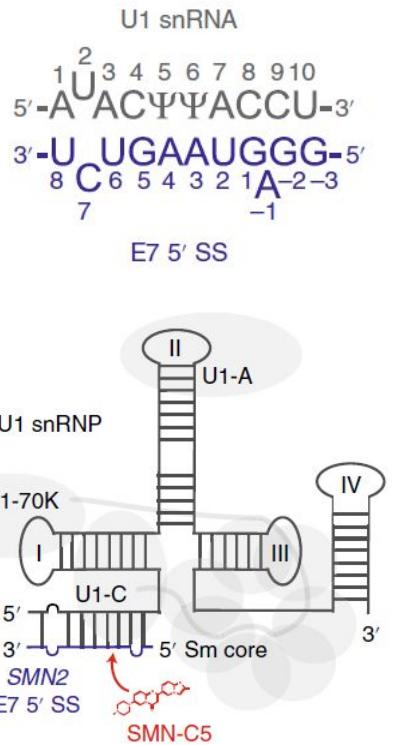
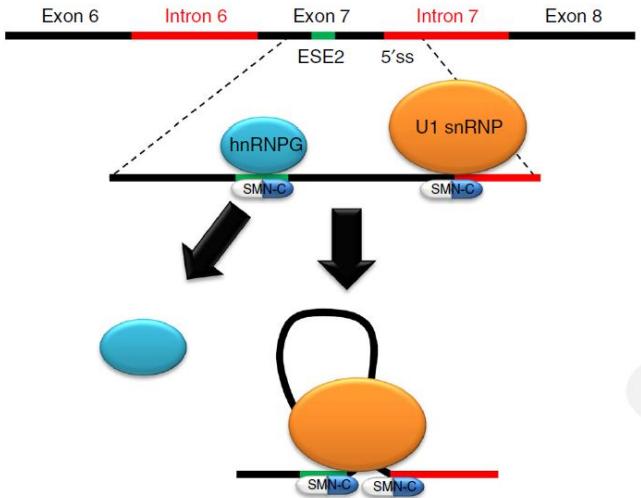
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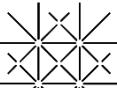


J



# Structural basis of specific splicing correction





# FIREFISH Part 1 Results

**Table 1.** Demographic and Clinical Characteristics of the Patients at Baseline.\*

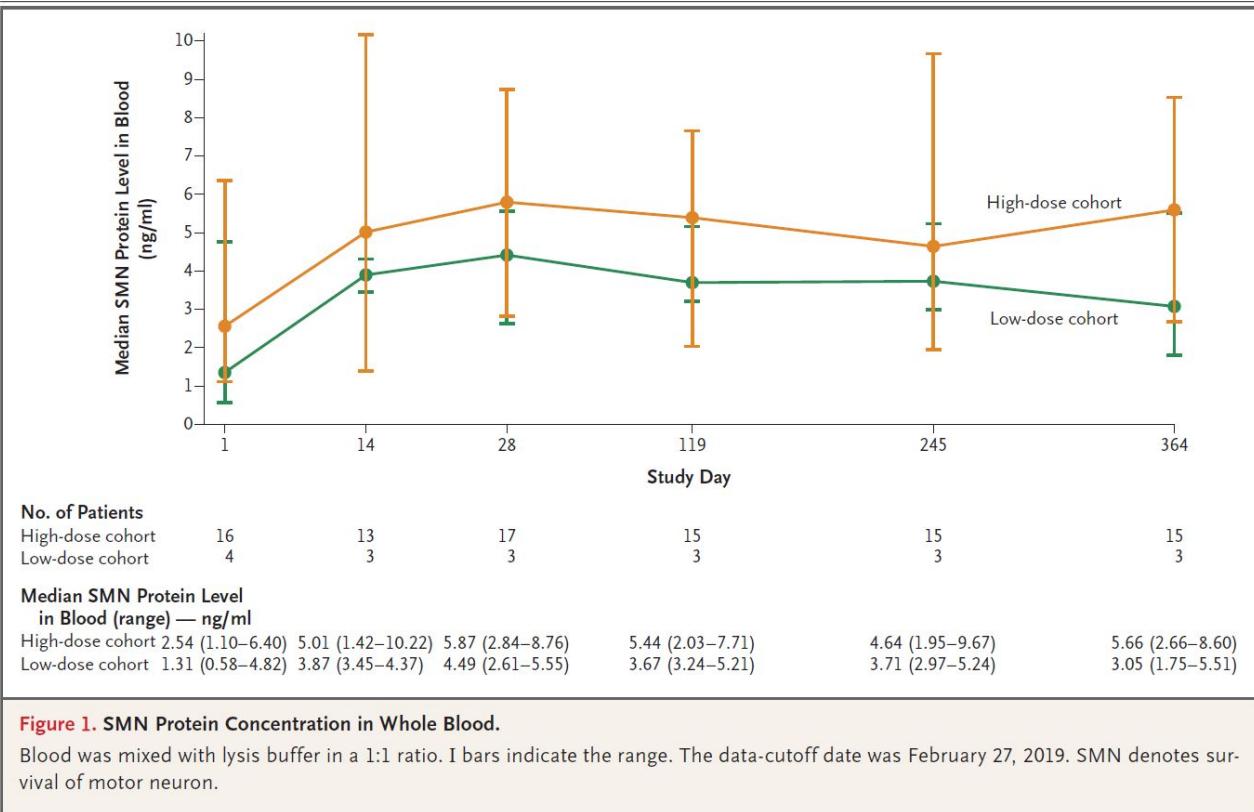
Characteristic	Low-Dose Cohort (N=4)	High-Dose Cohort (N=17)	All Infants (N=21)
Sex — no. (%)			
Female	4 (100)	11 (65)	15 (71)
Male	0	6 (35)	6 (29)
Median age (range) — mo			
At onset of symptoms	2.7 (2.0–3.0)	1.5 (0.9–3.0)	2.0 (0.9–3.0)
At diagnosis	3.3 (2.5–5.1)	3.0 (0.9–5.4)	3.0 (0.9–5.4)
At enrollment	6.9 (6.7–6.9)	6.3 (3.3–6.9)	6.7 (3.3–6.9)
Motor measures†			
Median CHOP-INTEND score (range)	23.5 (10–25)	24 (16–34)	24 (10–34)
Median HINE-2 score (range)	1 (0–3)	1 (0–2)	1 (0–3)
Respiratory support — no. (%)	0	5 (29)‡	5 (24)‡

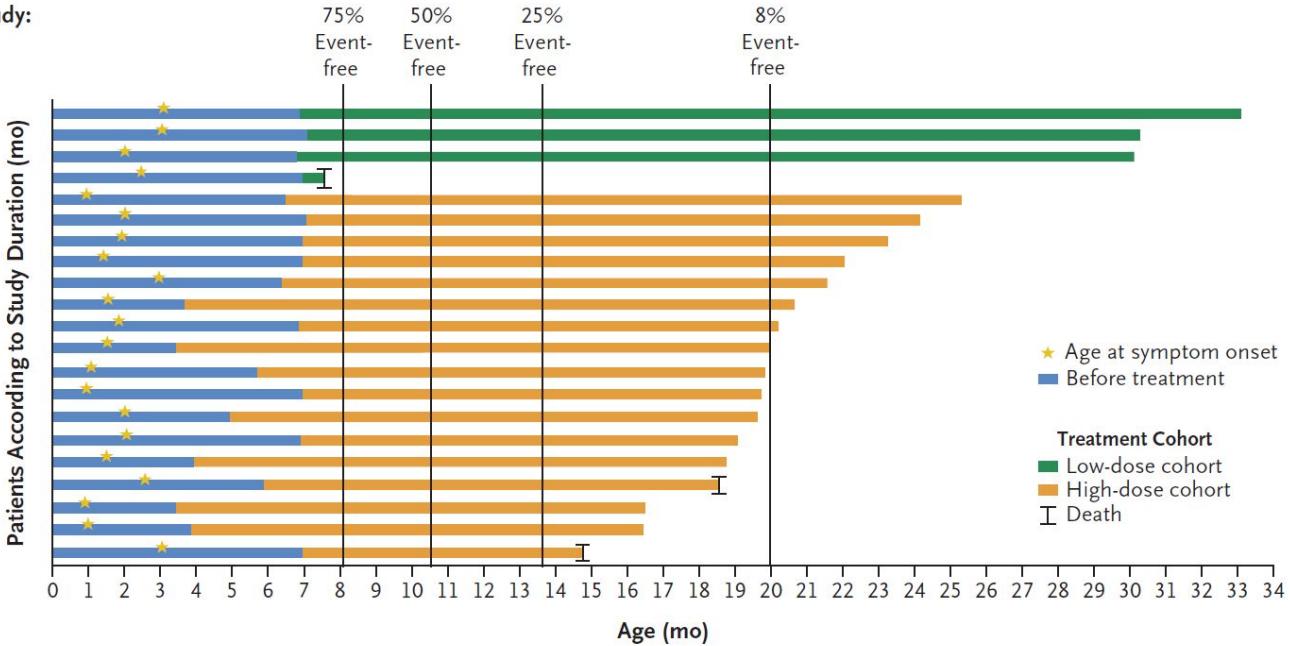
Note: Table 2 is not complete

**Table 2.** Adverse Events.\*

Event	Infants (N=21)
Total no. of adverse events	202
≥1 Adverse event — no. (%)	21 (100)
Total no. of serious adverse events	24
≥1 Serious adverse event — no. (%)	10 (48)
≥1 Adverse event of grade 3–5 — no. (%)	9 (43)
Serious adverse event with fatal outcome — no. (%)†	3 (14)
Most common adverse events — no. (%)‡	
Pyrexia	11 (52)
Upper respiratory tract infection	9 (43)
Diarrhea	6 (29)
Cough	5 (24)

# FIREFISH Part 1 Results

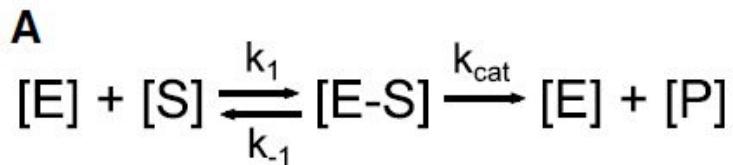


**Natural History Study:**


**Figure 2. Event-free Survival.**

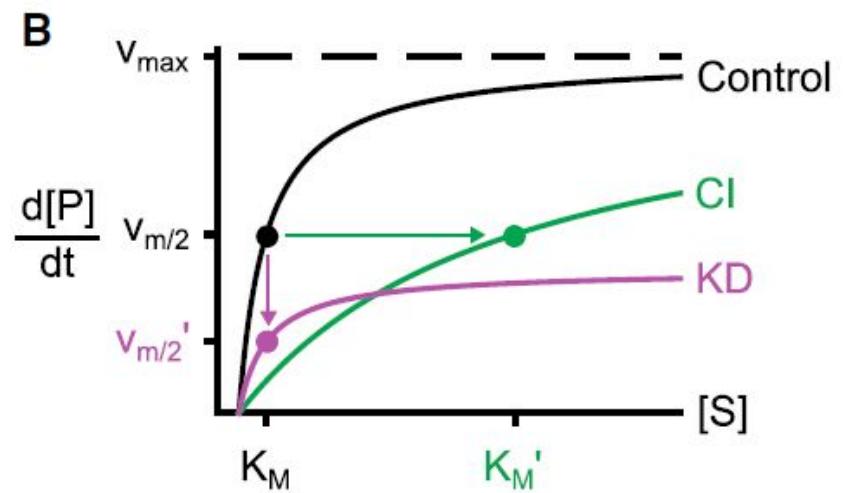
Event-free survival was defined as being alive and not receiving permanent ventilation (tracheostomy or ventilation [bilevel positive airway pressure] for  $\geq 16$  hours per day continuously for  $>3$  weeks or continuous intubation for  $>3$  weeks, in the absence of, or after the resolution of, an acute reversible event). The percentages of patients who were event-free in a previous natural history study of spinal muscular atrophy<sup>7</sup> are shown at the top of the graph for comparison. The median age at the combined outcome among patients in the previous study who had two copies of *SMN2* was 10.5 months (interquartile range, 8.1 to 13.6); event-free survival in that study was defined as being alive and not receiving noninvasive ventilation for 16 hours or more per day continuously for 2 or more weeks. The duration of our study was measured from the date of enrollment to the data-cutoff date. As of the data-cutoff date, three infants (one in the low-dose cohort and two in the high-dose cohort) had died; one additional infant in the high-dose cohort died after that date (Table S5).

# Enzymic and genetic inhibition have distinct impact on reaction dynamics



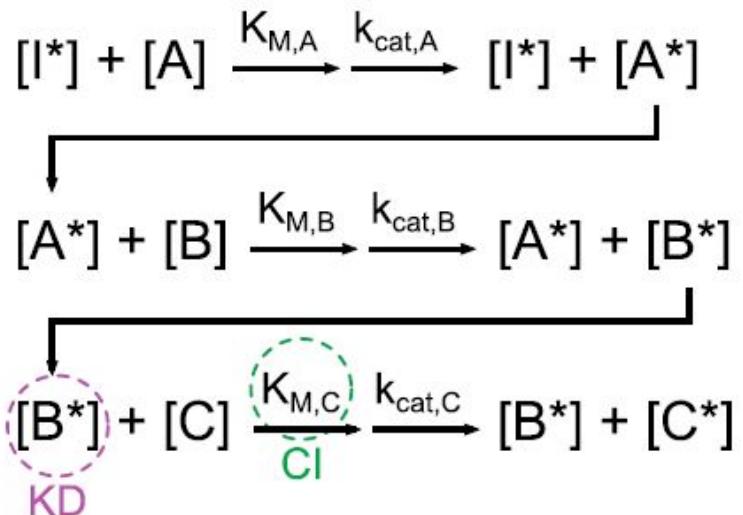
$$\frac{d[P]}{dt} = v_{\max} \frac{[S]}{[S] + K_M}$$

$$K_M = \frac{k_{-1} + k_{\text{cat}}}{k_1} \quad v_{\max} = k_{\text{cat}}[E]_0$$

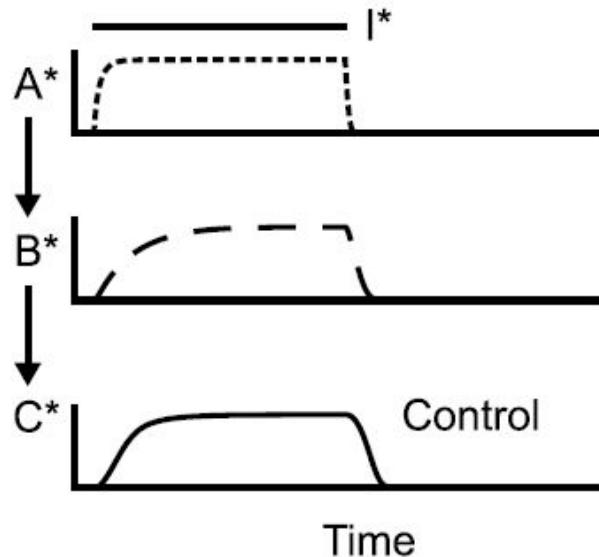


# A linear system of enzymatic reactions

C



D



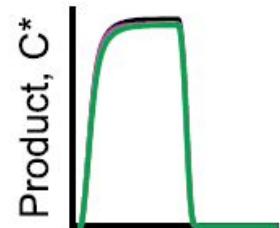
# Network architecture may differentiate effects of enzymatic and genetic inhibition

E

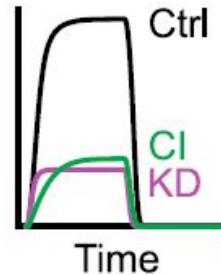
Linear



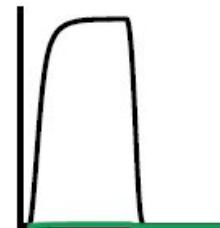
10% Inhibition



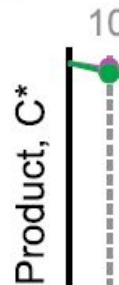
80% Inhibition



99% Inhibition



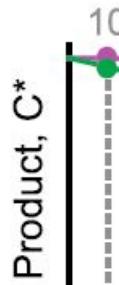
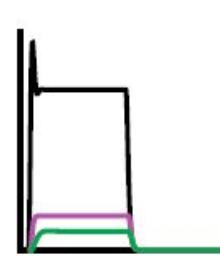
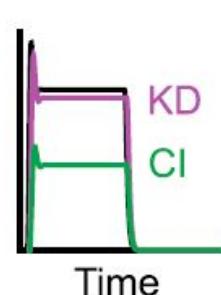
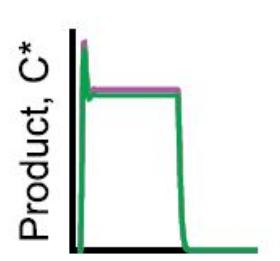
F



80 99

Inhibition of  $B \rightarrow C$

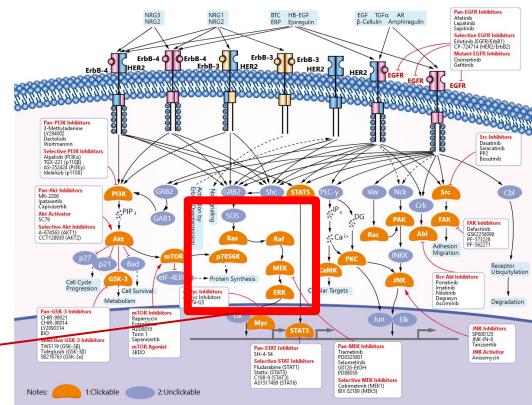
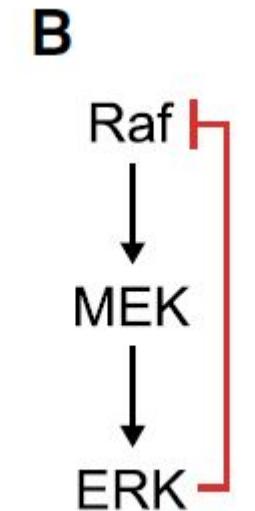
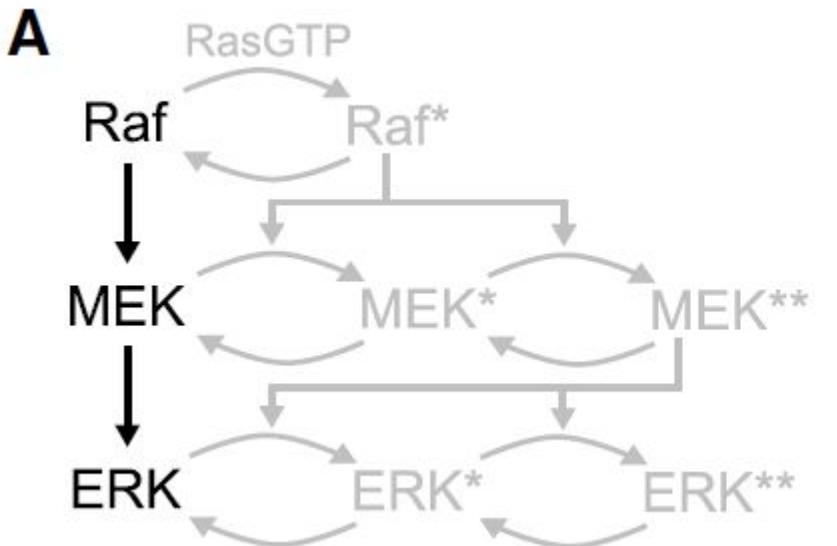
Feedback



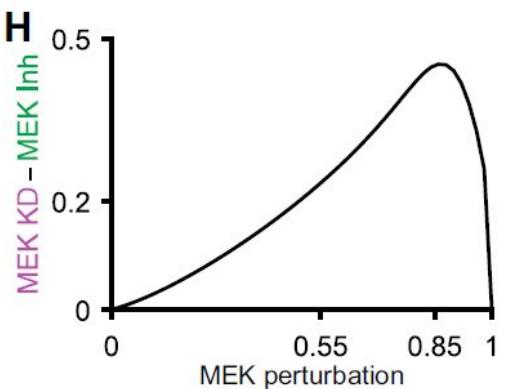
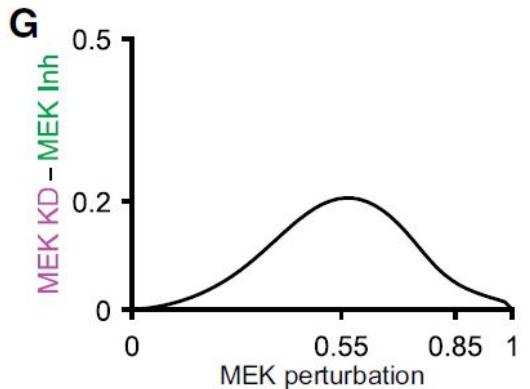
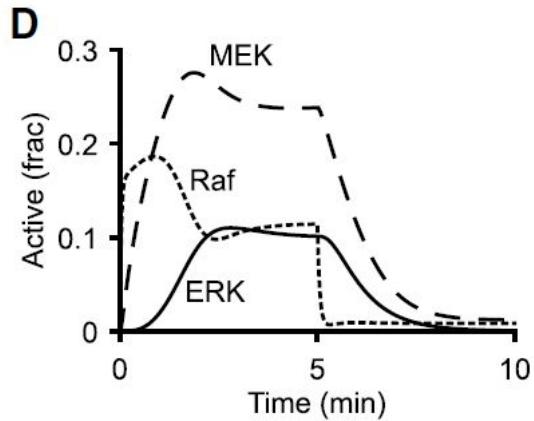
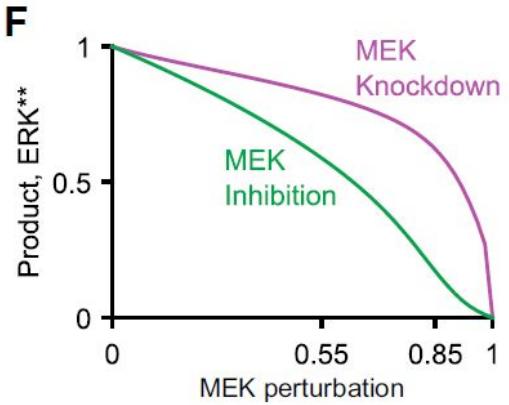
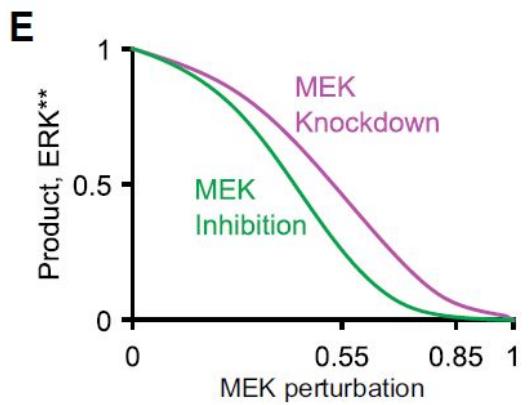
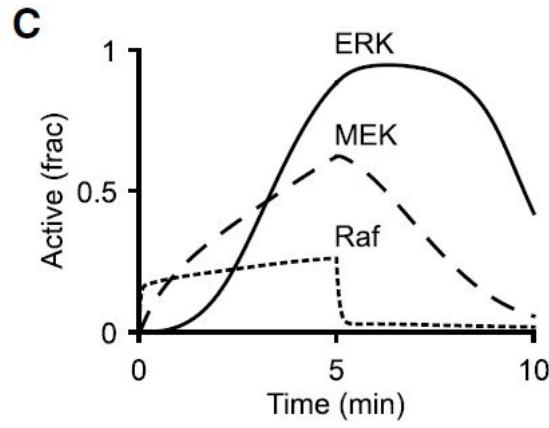
80 99

Inhibition of  $B \rightarrow C$

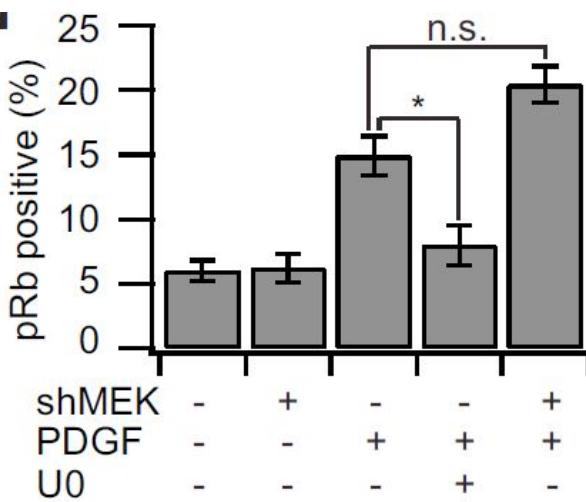
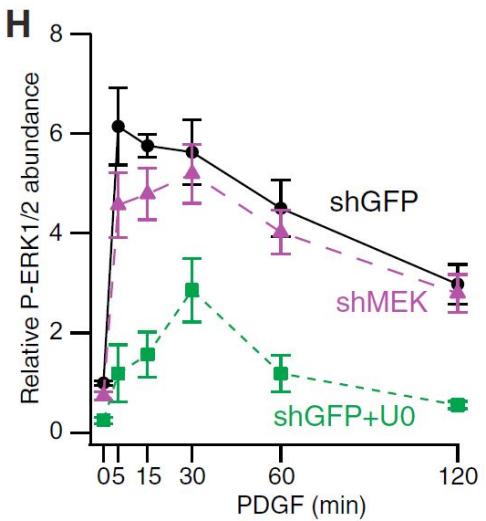
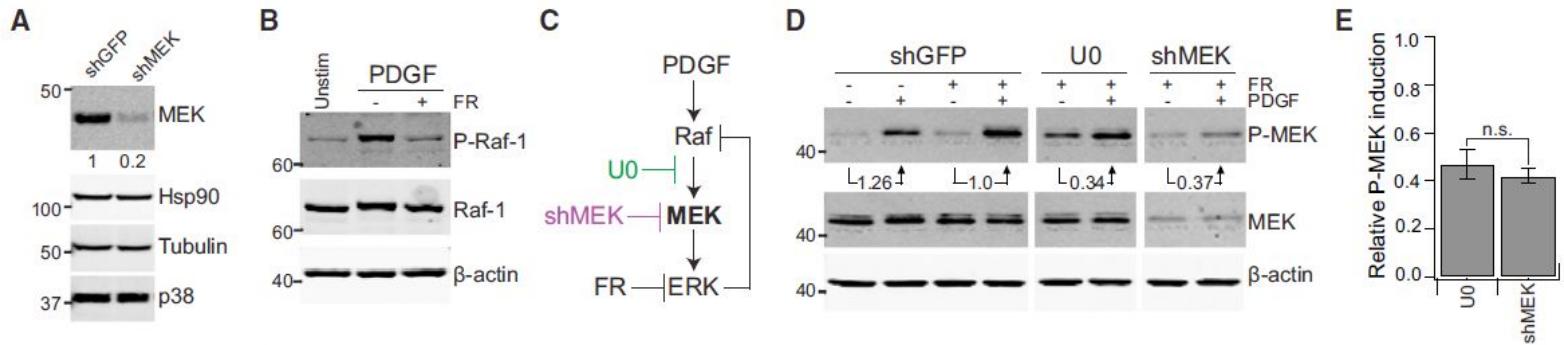
# The MAPK/ERK pathway downstream of EGFR signalling



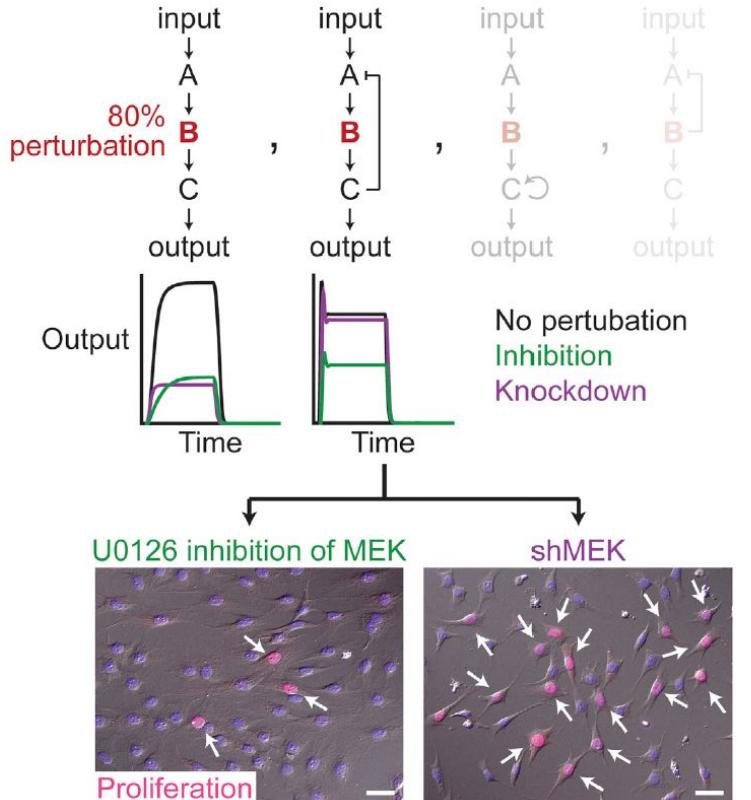
# A model predicts different effect of KD and CI



# Experimental data is in line with model prediction

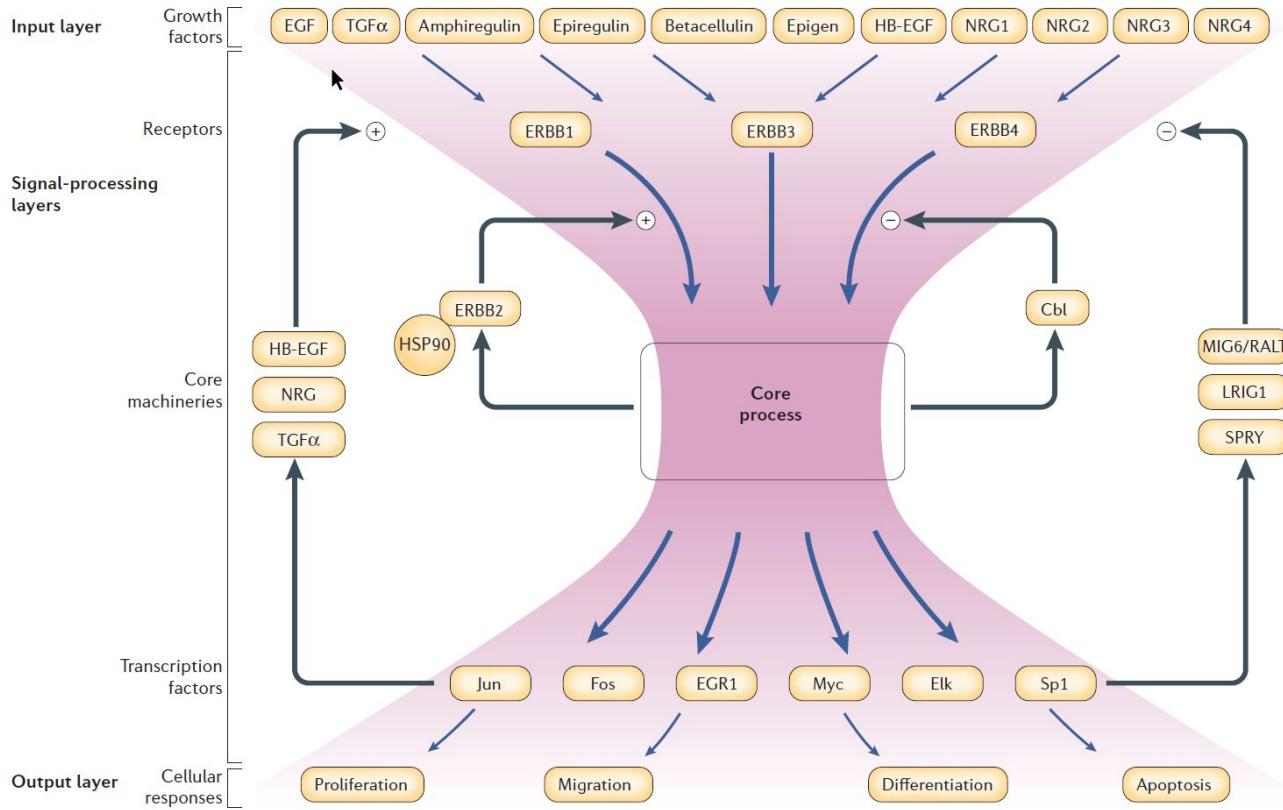


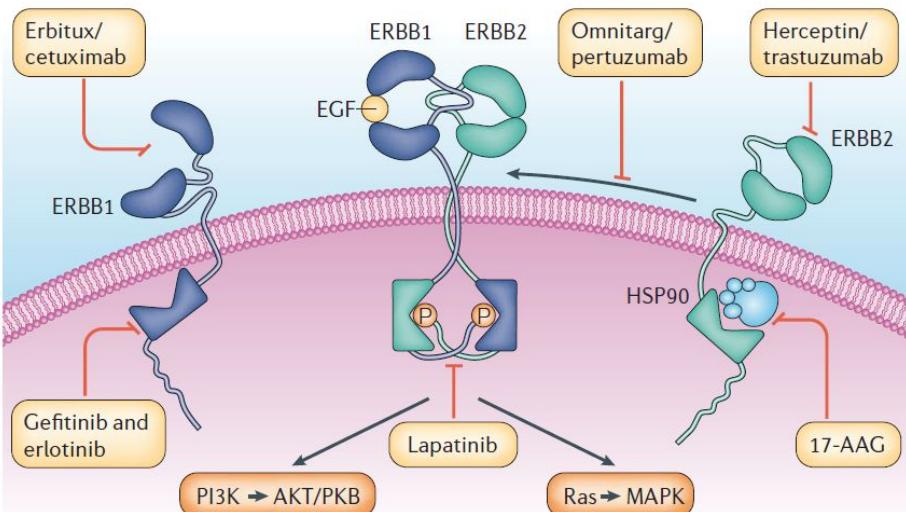
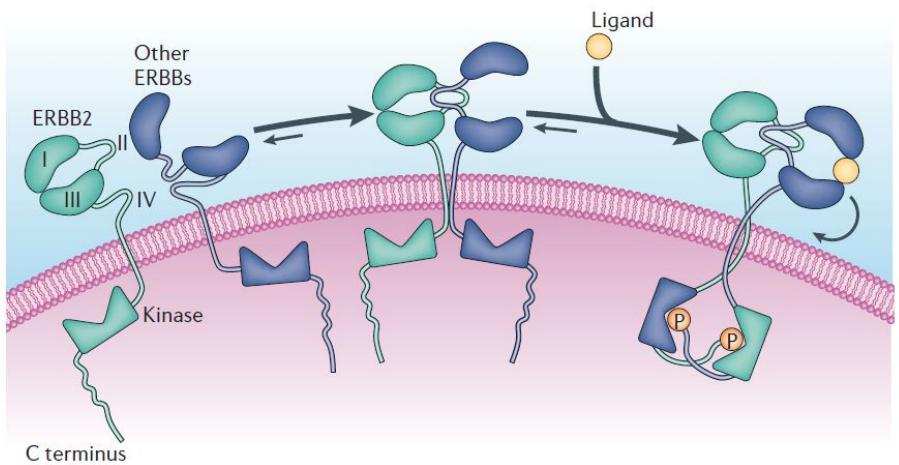
# Network architecture around an enzyme can affect our choice of modality

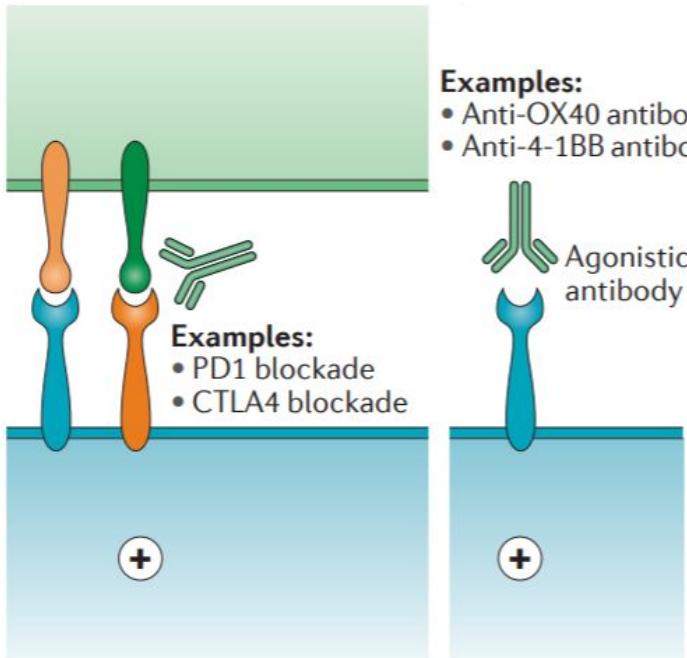
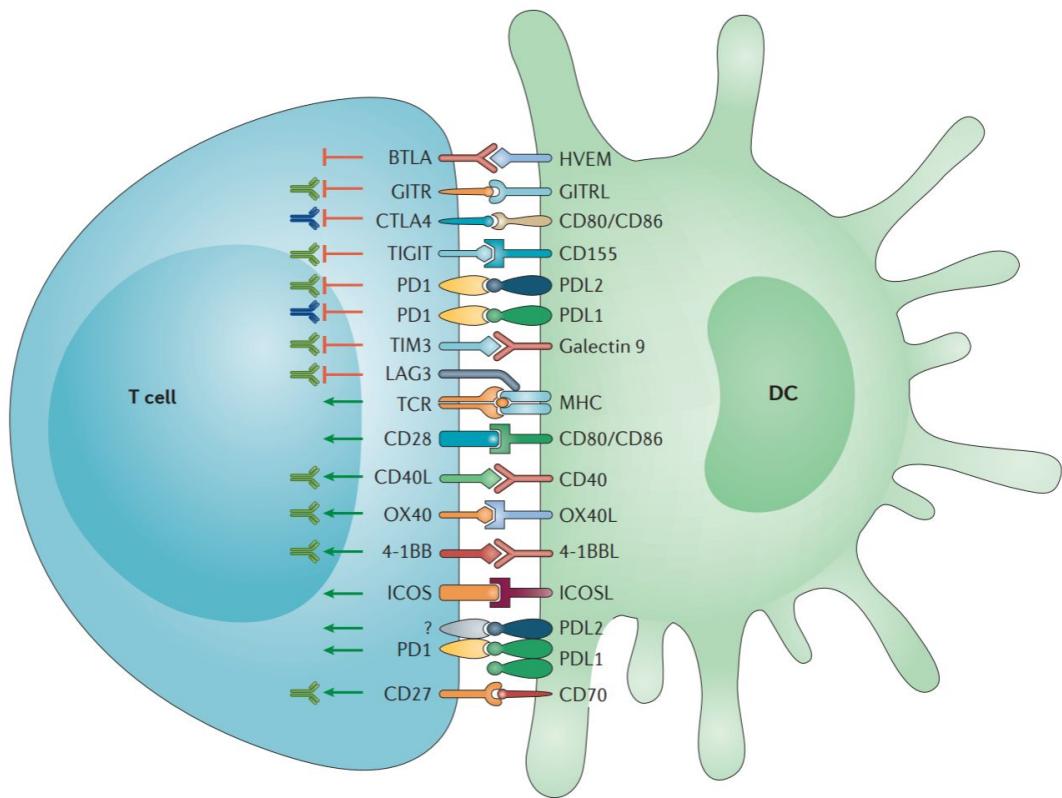


# Conclusions

- Given mechanistic understanding of biological processes underlying diseases, we can develop different modalities as therapeutics;
- Mathematical and computational biology (1) helps with molecule design, (2) reveals how drug candidate work and ranks them, and (3) contributes to modality selection;

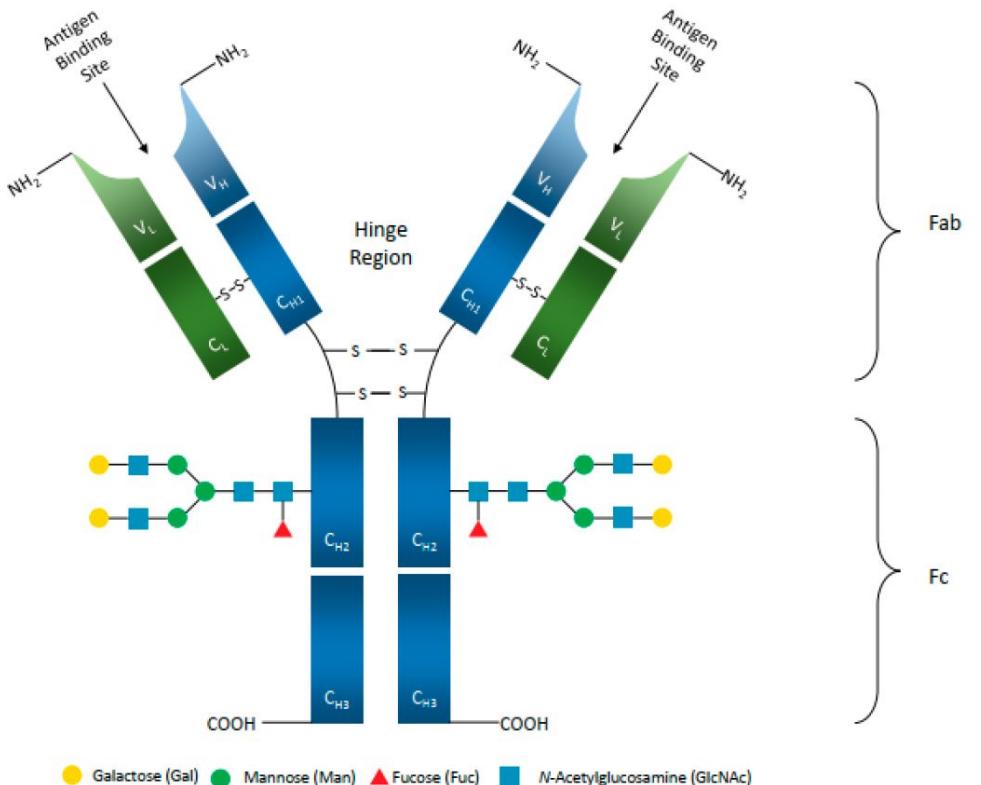




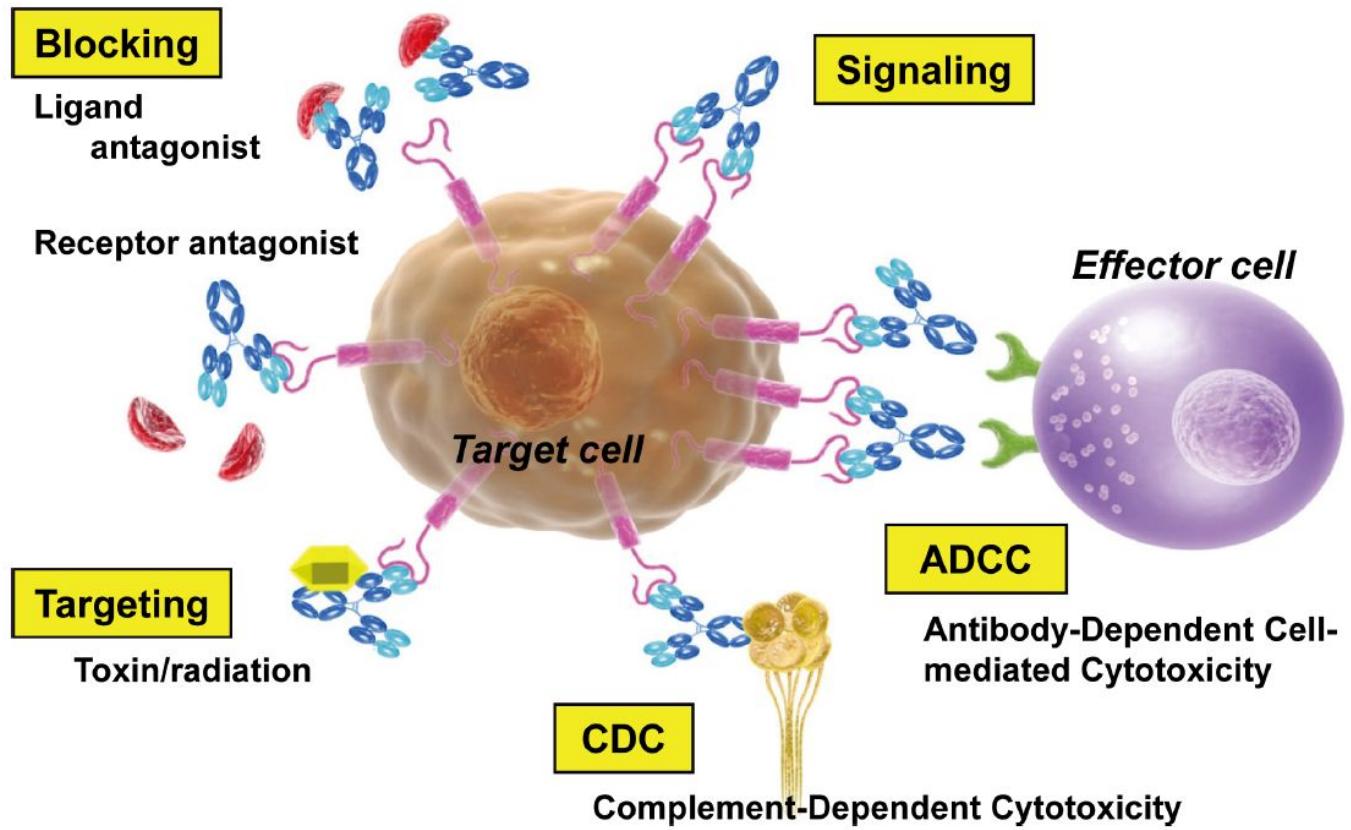


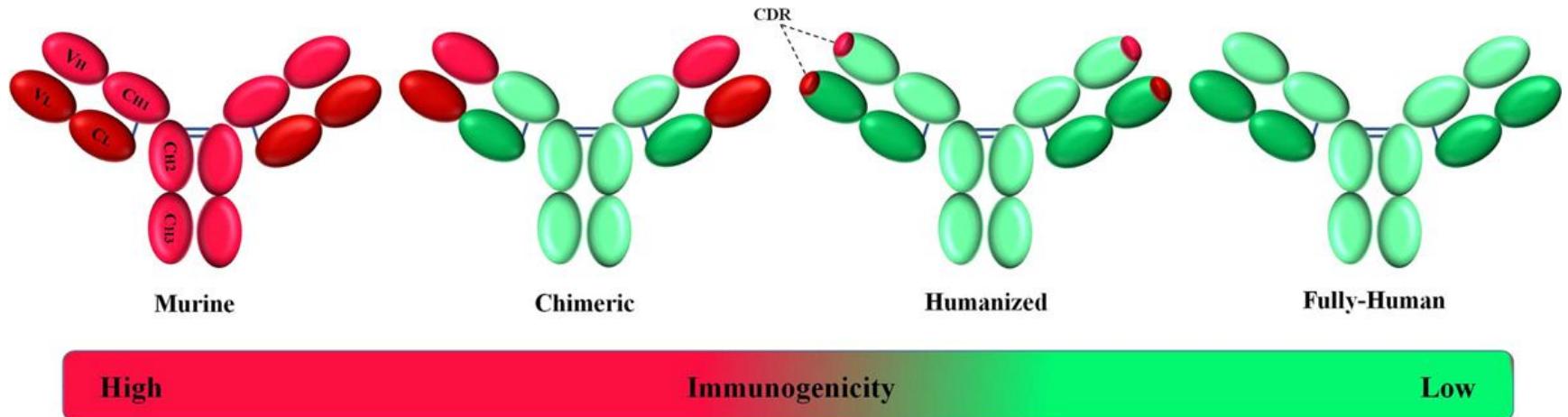
# Peptide and antibody

- Antibody design
- Immunogenicity



# Mechanisms of action of therapeutic antibodies

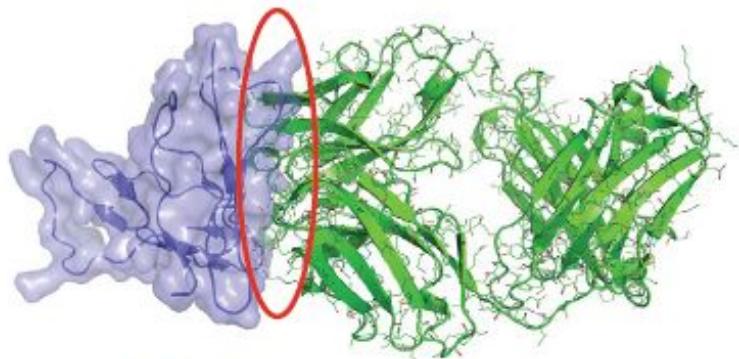




# Multispecific Drug Use or Target Interactions

**b** Conventional drug:

- Forms 1 drug–target interface
- Can act throughout body
- Only works if its binding to target alters function of target



IL-2R $\alpha$

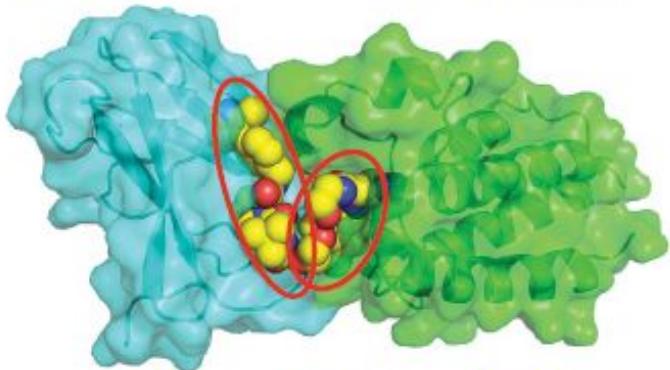
Basiliximab

**c** Obligate multispecific drug:

- Forms 2 or more drug–target interfaces

Class 1 'tetherbodies'  
 • Enrich drug at relevant site of action

Class 2 'matchmakers'  
 • Link drug to a biological effector



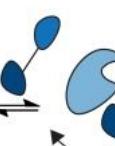
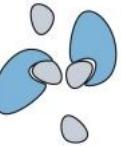
VHL

MZ1

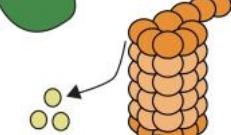
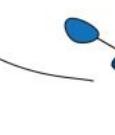
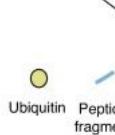
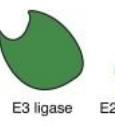
BRD4 $^{BD2}$

**(a) Occupancy-driven pharmacology**

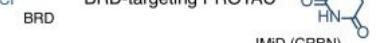
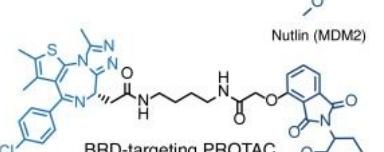
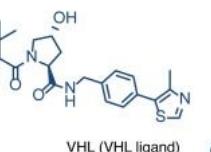
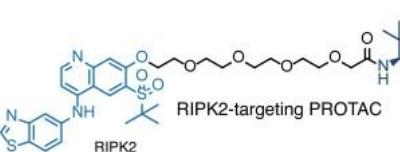
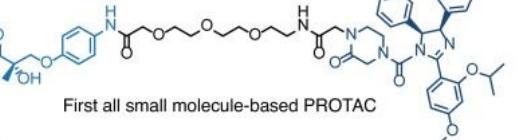
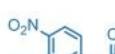
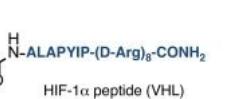
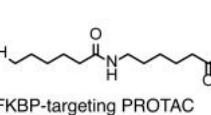
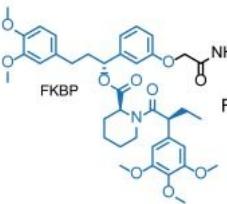
Protein function is modulated via inhibition


**(b) Event-driven pharmacology**

Protein function is modulated via PROTAC induced degradation



Catalytic process

**(c) PROteolysis TArgeting Chimera (PROTAC)**


# Offline Activities

- **Reading about RNA therapies:**
  - Levin, Arthur A. 2019. “Treating Disease at the RNA Level with Oligonucleotides.” *New England Journal of Medicine* 380 (1): 57–70.  
<https://doi.org/10.1056/NEJMra1705346>.
  - Oligonucleotides and their discontents:  
<https://blogs.sciencemag.org/pipeline/archives/2021/03/24/oligonucleotides-and-their-discontents>

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